Batch #4/7R
07/12/97

Serial Numb r: 09/147,919

CRF Processing Dat : 6/11/99
Edited by: _____
Verified by: _____ (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:

~~ENTERED~~
_____

☑ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☑ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: Th abov corrections must be communicated to the applicant in the first Offic Action. DO NOT send a copy of this form.

3/1/95

INPUT SET: S32192.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

```
 1                                     SEQUENCE LISTING
 2
 3      (1)     General Information:
 4
 5           (i) APPLICANT:
 6                (A) NAME: Bavarian Nordic AS
 7                (B) STREET: Naverland 2
 8                (C) CITY: Glostrup
 9                (E) COUNTRY: Denmark
10                (F) POSTAL CODE (ZIP): 2600
11                (G) TELEPHONE: oo45 43 438 444
12
13                (A) NAME: GSF Forschungszentrum fuer Umwelt und
14                          Gesundheit GmbH
15                (B) STREET: Ingolstaedter Landstr. 1
16                (C) CITY: Neuherberg
17                (E) COUNTRY: Germany
18                (F) POSTAL CODE (ZIP): 85758
19
20          (ii) TITLE OF INVENTION: Recombinant MVA Virus Expressing Dengue Virus
21                    Antigens, and The Use Thereof in Vaccines
22
23         (iii) NUMBER OF SEQUENCES: 8
24
25           (v) COMPUTER READABLE FORM:
26                (A) MEDIUM TYPE: Floppy disk
27                (B) COMPUTER: IBM PC compatible
28                (C) OPERATING SYSTEM: PC-DOS/MS-DOS
29                (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
30
31          (vi) CURRENT APPLICATION DATA:
32                (A) APPLICATION NUMBER:
33                (B) FILING DATE:
34
35         (vii) PRIOR APPLICATION DATA:
36                (A) APPLICATION NUMBER: DK 1035/96
37                (B) FILING DATE:
38
39      (2) INFORMATION FOR SEQ ID NO: 1:
40
41           (i) SEQUENCE CHARACTERISTICS:
42                (A) LENGTH: 33 base pairs
43                (B) TYPE: nucleic acid
44                (C) STRANDEDNESS: single
45                (D) TOPOLOGY: linear
46
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The following lines are marked with --> in the margin:
5, 6, 7, 8, 9, 10, 11, 13, 15, 16, 17, 18

ENTERED

RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/147,919*

INPUT SET: S32192.raw

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47          (ii) MOLECULE TYPE: other nucleic acid
48                (A) DESCRIPTION:   /desc = "primer"
49
50         (iii) HYPOTHETICAL: YES
51
52          (iv) ANTI-SENSE: NO
53
54
55
56
57          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
58
59  CAGCAGGGTA CCCTCATCGT ACAGGACGTT CTC                                  33
60
61  (2) INFORMATION FOR SEQ ID NO: 2:
62
63           (i) SEQUENCE CHARACTERISTICS:
64                (A) LENGTH: 42 base pairs
65                (B) TYPE: nucleic acid
66                (C) STRANDEDNESS: single
67                (D) TOPOLOGY: linear
68
69          (ii) MOLECULE TYPE: other nucleic acid
70                (A) DESCRIPTION:   /desc = "primer"
71
72
73
74
75          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
76
77  CAGCAGCCCG GGTATTCGAT GATTATTTTT AACAAAATAA CA                        42
78
79  (2) INFORMATION FOR SEQ ID NO: 3:
80
81           (i) SEQUENCE CHARACTERISTICS:
82                (A) LENGTH: 36 base pairs
83                (B) TYPE: nucleic acid
84                (C) STRANDEDNESS: single
85                (D) TOPOLOGY: linear
86
87          (ii) MOLECULE TYPE: other nucleic acid
88                (A) DESCRIPTION:   /desc = "primer"
89
90         (iii) HYPOTHETICAL: YES
91
92          (iv) ANTI-SENSE: NO
93
94
95
96
97          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
98
99  CAGCAGCTGC AGGAATCATC CATTCCACTG AATAGC                               36
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PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/147,919*

DATE: 06/14/1999
TIME: 16:45:39

INPUT SET: S32192.raw

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100
101    (2) INFORMATION FOR SEQ ID NO: 4:
102
103        (i) SEQUENCE CHARACTERISTICS:
104            (A) LENGTH: 36 base pairs
105            (B) TYPE: nucleic acid
106            (C) STRANDEDNESS: single
107            (D) TOPOLOGY: linear
108
109       (ii) MOLECULE TYPE: other nucleic acid
110            (A) DESCRIPTION:   /desc = "primer"
111
112
113
114
115       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
116
117  CAGCAGGCAT GCCGACGAAC AAGGAACTGT AGCAGA                              36
118
119    (2) INFORMATION FOR SEQ ID NO: 5:
120
121        (i) SEQUENCE CHARACTERISTICS:
122            (A) LENGTH: 33 base pairs
123            (B) TYPE: nucleic acid
124            (C) STRANDEDNESS: single
125            (D) TOPOLOGY: linear
126
127       (ii) MOLECULE TYPE: other nucleic acid
128            (A) DESCRIPTION:   /desc = "primer"
129
130
131
132
133       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
134
135  CAGCAGGTCG ACCCCGACCG CCTTACTGCC GCC                                 33
136
137    (2) INFORMATION FOR SEQ ID NO: 6:
138
139        (i) SEQUENCE CHARACTERISTICS:
140            (A) LENGTH: 33 base pairs
141            (B) TYPE: nucleic acid
142            (C) STRANDEDNESS: single
143            (D) TOPOLOGY: linear
144
145       (ii) MOLECULE TYPE: other nucleic acid
146            (A) DESCRIPTION:   /desc = "primer"
147
148
149
150
151       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
152
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INPUT SET: S32192.raw

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153   GGGGGGCTGC AGATGGTAGC GACCGGCGCT CAG                           33
154
155   (2) INFORMATION FOR SEQ ID NO: 7:
156
157       (i) SEQUENCE CHARACTERISTICS:
158           (A) LENGTH: 36 base pairs
159           (B) TYPE: nucleic acid
160           (C) STRANDEDNESS: single
161           (D) TOPOLOGY: linear
162
163       (ii) MOLECULE TYPE: other nucleic acid
164           (A) DESCRIPTION:   /desc = "primer"
165
166
167
168
169       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
170
171   CAGCAGCCCG GGATGGCAGC AATCCTGGCA TACACC                        36
172
173   (2) INFORMATION FOR SEQ ID NO: 8:
174
175       (i) SEQUENCE CHARACTERISTICS:
176           (A) LENGTH: 39 base pairs
177           (B) TYPE: nucleic acid
178           (C) STRANDEDNESS: single
179           (D) TOPOLOGY: linear
180
181       (ii) MOLECULE TYPE: other nucleic acid
182           (A) DESCRIPTION:   /desc = "primer"
183
184
185
186
187       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
188
189   CAGCAGCCCG GGTCACTGTA GTTTGTCCAT CCTCAGCCT                      39
190
```

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION *US/09/147,919*

INPUT SET: S32192.raw

Line	Error	Original Text
5	Mandatory Value Not Present	(i) APPLICANT:
6	Unknown or Misplaced Identifier	(A) NAME: Bavarian Nordic AS
7	Unknown or Misplaced Identifier	(B) STREET: Naverland 2
8	Unknown or Misplaced Identifier	(C) CITY: Glostrup
9	Unknown or Misplaced Identifier	(E) COUNTRY: Denmark
10	Unknown or Misplaced Identifier	(F) POSTAL CODE (ZIP): 2600
11	Unknown or Misplaced Identifier	(G) TELEPHONE: oo45 43 438 444
13	Unknown or Misplaced Identifier	(A) NAME: GSF Forschungszentrum fuer Umwelt und
15	Unknown or Misplaced Identifier	(B) STREET: Ingolstaedter Landstr. 1
16	Unknown or Misplaced Identifier	(C) CITY: Neuherberg
17	Unknown or Misplaced Identifier	(E) COUNTRY: Germany
18	Unknown or Misplaced Identifier	(F) POSTAL CODE (ZIP): 85758

1643 BATCH

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/147,919*

DATE: 06/11/1999
TIME: 10:23:23

INPUT SET: S32192.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

```
 1                              SEQUENCE LISTING
 2
 3    (1)    General Information:
 4
 5          (i) APPLICANT:
 6              (A) NAME: Bavarian Nordic AS
 7              (B) STREET: Naverland 2
 8              (C) CITY: Glostrup
 9              (E) COUNTRY: Denmark
10              (F) POSTAL CODE (ZIP): 2600
11              (G) TELEPHONE: oo45 43 438 444
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13              (A) NAME: GSF Forschungszentrum fuer Umwelt und
14                        Gesundheit GmbH
15              (B) STREET: Ingolstaedter Landstr. 1
16              (C) CITY: Neuherberg
17              (E) COUNTRY: Germany
18              (F) POSTAL CODE (ZIP): 85758
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20         (ii) TITLE OF INVENTION: Recombinant MVA Virus Expressing Dengue Virus
21                 Antigens, and The Use Thereof in Vaccines
22
23        (iii) NUMBER OF SEQUENCES: 8
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25         (iv) COMPUTER READABLE FORM:
26              (A) MEDIUM TYPE: Floppy disk
27              (B) COMPUTER: IBM PC compatible
28              (C) OPERATING SYSTEM: PC-DOS/MS-DOS
29              (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
30
31         (vi) CURRENT APPLICATION DATA:
32              (A) APPLICATION NUMBER: DK 1035/96
33
34    (2) INFORMATION FOR SEQ ID NO: 1:
35
36         (i) SEQUENCE CHARACTERISTICS:
37              (A) LENGTH: 33 base pairs
38              (B) TYPE: nucleic acid
39              (C) STRANDEDNESS: single
40              (D) TOPOLOGY: linear
41
42        (ii) MOLECULE TYPE: other nucleic acid
43              (A) DESCRIPTION:   /desc = "primer"
44
45       (iii) HYPOTHETICAL: YES
46
```

Does Not Comply
Corrected Diskette Needed

(vii) PRIOR APP DATA!

```
47        (iv) ANTI-SENSE: NO
48
49
50
51
52        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
53
54   CAGCAGGGTA CCCTCATCGT ACAGGACGTT CTC                          33
55
56   (2) INFORMATION FOR SEQ ID NO: 2:
57
58        (i) SEQUENCE CHARACTERISTICS:
59             (A) LENGTH: 42 base pairs
60             (B) TYPE: nucleic acid
61             (C) STRANDEDNESS: single
62             (D) TOPOLOGY: linear
63
64        (ii) MOLECULE TYPE: other nucleic acid
65             (A) DESCRIPTION:   /desc = "primer"
66
67
68
69
70        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
71
72   CAGCAGCCCG GGTATTCGAT GATTATTTTT AACAAAATAA CA                 42
73
74   (2) INFORMATION FOR SEQ ID NO: 3:
75
76        (i) SEQUENCE CHARACTERISTICS:
77             (A) LENGTH: 36 base pairs
78             (B) TYPE: nucleic acid
79             (C) STRANDEDNESS: single
80             (D) TOPOLOGY: linear
81
82        (ii) MOLECULE TYPE: other nucleic acid
83             (A) DESCRIPTION:   /desc = "primer"
84
85       (iii) HYPOTHETICAL: YES
86
87        (iv) ANTI-SENSE: NO
88
89
90
91
92        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
93
94   CAGCAGCTGC AGGAATCATC CATTCCACTG AATAGC                       36
95
96   (2) INFORMATION FOR SEQ ID NO: 4:
97
98        (i) SEQUENCE CHARACTERISTICS:
99             (A) LENGTH: 36 base pairs
```

```
100            (B) TYPE: nucleic acid
101            (C) STRANDEDNESS: single
102            (D) TOPOLOGY: linear
103
104       (ii) MOLECULE TYPE: other nucleic acid
105            (A) DESCRIPTION:   /desc = "primer"
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109
110       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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112 CAGCAGGCAT GCCGACGAAC AAGGAACTGT AGCAGA                           36
113
114 (2) INFORMATION FOR SEQ ID NO: 5:
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116        (i) SEQUENCE CHARACTERISTICS:
117            (A) LENGTH: 33 base pairs
118            (B) TYPE: nucleic acid
119            (C) STRANDEDNESS: single
120            (D) TOPOLOGY: linear
121
122       (ii) MOLECULE TYPE: other nucleic acid
123            (A) DESCRIPTION:   /desc = "primer"
124
125
126
127
128       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
129
130 CAGCAGGTCG ACCCCGACCG CCTTACTGCC GCC                              33
131
132 (2) INFORMATION FOR SEQ ID NO: 6:
133
134        (i) SEQUENCE CHARACTERISTICS:
135            (A) LENGTH: 33 base pairs
136            (B) TYPE: nucleic acid
137            (C) STRANDEDNESS: single
138            (D) TOPOLOGY: linear
139
140       (ii) MOLECULE TYPE: other nucleic acid
141            (A) DESCRIPTION:   /desc = "primer"
142
143
144
145
146       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
147
148 GGGGGGCTGC AGATGGTAGC GACCGGCGCT CAG                              33
149
150 (2) INFORMATION FOR SEQ ID NO: 7:
151
152        (i) SEQUENCE CHARACTERISTICS:
```

```
153                (A) LENGTH: 36 base pairs
154                (B) TYPE: nucleic acid
155                (C) STRANDEDNESS: single
156                (D) TOPOLOGY: linear
157
158          (ii) MOLECULE TYPE: other nucleic acid
159                (A) DESCRIPTION:   /desc = "primer"
160
161
162
163
164          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
165
166   CAGCAGCCCG GGATGGCAGC AATCCTGGCA TACACC                        36
167
168   (2) INFORMATION FOR SEQ ID NO: 8:
169
170           (i) SEQUENCE CHARACTERISTICS:
171                (A) LENGTH: 39 base pairs
172                (B) TYPE: nucleic acid
173                (C) STRANDEDNESS: single
174                (D) TOPOLOGY: linear
175
176          (ii) MOLECULE TYPE: other nucleic acid
177                (A) DESCRIPTION:   /desc = "primer"
178
179
180
181
182          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
183
184   CAGCAGCCCG GGTCACTGTA GTTTGTCCAT CCTCAGCCT                     39
185
```

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION *US/09/147,919*

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Line	Error	Original Text
5	Mandatory Value Not Present	(i) APPLICANT:
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10	Unknown or Misplaced Identifier	(F) POSTAL CODE (ZIP): 2600
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16	Unknown or Misplaced Identifier	(C) CITY: Neuherberg
17	Unknown or Misplaced Identifier	(E) COUNTRY: Germany
18	Unknown or Misplaced Identifier	(F) POSTAL CODE (ZIP): 85758